PERFORMANCE REPORT

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Title: **DNA Computing**

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Form Approved REPORT DOCUMENTATION PAGE OMB No. 0704-0188 Public reporting burden for this collection of information is estimated to average 1 hour per response, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing this collection of information. Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden to Department of Defense, Washington Headquarters Services, Directorate for Information Operations and Reports (0704-0188), 1215 Jefferson Davis Highway, Suite 1204, Arlington, VA 22202-4302. Respondents should be aware that notwithstanding any other provision of law, no person shall be subject to any penalty for failing to comply with a collection of information if it does not display a currently valid OMB control number. PLEASE DO NOT RETURN YOUR FORM TO THE ABOVE ADDRESS. 1. REPORT DATE (DD-MM-YYYY) 3. DATES COVERED (From - To) 2. REPORT TYPE 21-03-2008 Final 4. TITLE AND SUBTITLE 5a. CONTRACT NUMBER DNA Computing **5b. GRANT NUMBER** FA9550-05-1-0424 **5c. PROGRAM ELEMENT NUMBER** 6. AUTHOR(S) 5d. PROJECT NUMBER 5e. TASK NUMBER Wayne Frasch 5f. WORK UNIT NUMBER 7. PERFORMING ORGANIZATION NAME(S) AND ADDRESS(ES) 8. PERFORMING ORGANIZATION REPORT NUMBER Arizona State University, School of Life Sciences, P.O. Box 874501, Tempe, AZ 85287-4501 9. SPONSORING / MONITORING AGENCY NAME(S) AND ADDRESS(ES) 10. SPONSOR/MONITOR'S ACRONYM(S) Air Force Office of Scientific Research **AFOSR** 875 N. Randolph St Arlington, VA 22203 11. SPONSOR/MONITOR'S REPORT NUMBER(S) AFRL-SR-AR-TR-08-0241 12. DISTRIBUTION / AVAILABILITY STATEMENT Approved for public release; distribution is unlimited 13. SUPPLEMENTARY NOTES 14. ABSTRACT A microscopy system was acquired for molecular detection using molecular motor-driven nanodevices. This state of the art system incorporates a motorized stage that can move to successive adjacent fields of view, and stitch the digital color photos for analysis. The detection system utilizes the molecular motor attached to a slide and a gold nanorod that is visible by darkfield microscopy. Assembly of each nanodevice is dependent upon the presence of one molecule of target that bridges between motor and nanorod. Zeptomole sensitivity of target DNA is achieved without PCR or other means of amplification via molecular motor-dependent rotation. We have now demonstrated simultaneous, multiplexed detection of sequence specific DNA, RNA, and proteins unique to MRSA on the same platform with three separate colors of gold nanorods each having a different functionalization. We have also demonstrated the ability to differentiate between Staphylococcus aureus and MRSA as well as detection of target from crude cell lysate. We have written algorithms that can analyze the stitched digital photos of fields of view, correctly identify red, green, and yellow nanorods used to detect DNA, RNA, and proteins, respectively, then rapidly quantify the amount of each target present. Assembly of a commercial prototype of the detection device is underway. 15. SUBJECT TERMS

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TASK A: DNA COMPUTING TO SOLVE OPTIMIZATION PROBLEMS

Funded by AFOSR and DARPA

DNA hybridization was used to make probabilistic computations to identify the optimal path for fully connected asymmetric Traveling Salesman Problems in four steps: (1) Answer generation; (2) Answer sorting; (3) Answer Readout; (4) Optimal Answer Identification. The DNA concentration of starting sequences was controlled to allow the entire answer space to be sampled, with the constraint that the probability of forming a particular answer sequence was directly proportional to its optimality. We first used this procedure to solve an asymmetric 10 city TSP with small concentration differences of starting sequences, which tested the ability to find optimal paths among many closely suboptimal answers out of ~3.3 million possible answers.

We now report the ability to identify the optimal answer of a fully asymmetric 15 City TSP for which ~1.3x10¹² possible solutions exist. Starting DNA concentrations were used that clearly favored formation of one optimal answer, in contrast to the 10 city TSP problem. The high abundance of a subset of DNA to form the optimal answers relative to suboptimal answers increased the difficulty of optimal answer sorting and readout due to problems in PCR signal amplification. This was averted by generating enough initial answer without PCR amplification to allow purification of correct answers. Under these conditions it was possible to identify the optimal answer to the 15 City TSP by inspection directly from raw data, and is the largest problem ever solved by DNA computing. An answer pool for a 20 City TSP was also successfully generated, and answer-sorting completed. Answer analysis is underway.

We have created a novel "double-padlock" probe to increase the speed and accuracy during answer determination. This will allow the use of RT-PCR to rapidly quantitate amounts of city pairings formed in the answer sequences. Construction has also begun of 3D DNA structures for DNA computations. These structures are being designed to: (1) eliminate incorrect answer formation that consumes valuable DNA starting material, and (2) use thermodynamic stability to drive optimal answer generation in order to minimize formation of suboptimal answers. We have used a supercomputing cluster to optimize the thermodynamic stability of proposed 3D DNA structures prior to synthesis of components and assembly.

TASK B: MICROSCOPY SYSTEM FOR MOLECULAR DETECTION USING NANODEVICES

Funded by AFOSR, DARPA, and DURIP

A microscopy system was acquired for molecular detection using molecular motor-driven nanodevices. This state of the art system incorporates a motorized stage that can move to successive adjacent fields of view, and stitch the digital color photos for analysis. The detection system utilizes the molecular motor attached to a slide and a gold nanorod that is visible by dark-field microscopy. Assembly of each nanodevice is dependent upon the presence of one molecule of target that bridges between motor and nanorod. Zeptomole sensitivity of target DNA is achieved without PCR or other means of amplification via molecular motor-dependent rotation. We have now demonstrated simultaneous, multiplexed detection of sequence specific DNA, RNA, and proteins unique to MRSA on the same platform with three separate colors of gold nanorods each having a different functionalization. We have also demonstrated the ability to differentiate between *Staphylococcus aureus* and MRSA as well as detection of target from crude cell lysate. We have written algorithms that can analyze the stitched digital photos of fields of view, correctly identify red, green, and yellow nanorods used to detect DNA, RNA, and proteins, respectively, then rapidly quantify the amount of each target present. Assembly of a commercial prototype of the detection device is underway.

Publications during past year:

- 1. Spetzler, D., Xiong, F., and Frasch, W.D. (2007) "Probabilistic DNA Computing Solution to a Fully Connected 10-City Asymmetric Traveling Salesman Problem" Proc. DNA 13, 9-18.
- 2. Xiong, F., Spetzler, and Frasch, W. D. (2007) "Elimination of Secondary Structures for DNA Computing", Proc. DNA 13, 241-249.
- 3. Spetzler, D., York, J., Dobbin, C., Martin, J., Xiong, F., Ishmukhametov, R., Day, L., Yu, J., Kang, H., Porter, K., Hornung, T., and Frasch, W.D. (2007) "Recent Developments of Biomolecular Motors as On-Chip Devices using Single Molecule Techniques", Lab. Chip 7, 1633-1643
- 4. Spetzler, D., Xiong, F., and Frasch, W. D. (2008) "Heuristic solution to a 10-City Traveling Salesman Problem Using Probabilistic DNA Computing", LNCS 4848, 152-160.
- York, J., Spetzler, D., Hornung, T., Ishmukhametov, R., Martin, J., and Frasch, W.D. (2007) "Abundance of Escherichia coli F1-ATPase Molecules Observed to Rotate via Single-Molecule Microscopy with Gold Nanorod Probes", J. Bioenergetics and Biomembranes, in press.
- 6. Spetzler, D., York, J., Martin. J., Ishmukhametov, R., and Frasch, W. D. "Microsecond Resolution of Enzymatic Movement Using Dark-Field Microscopy", Methods, in press. 7. York, J., Spetzler, D., Xiong, F., and Frasch, W.D. (2008) "Single-Molecule Detection of DNA via Sequence-Specific Links between F1-ATPase Motors and Gold Nanorod Sensors", Lab Chip., in press.

Patents during past year:

- 1. Frasch, W.D., Spetzler, D. and York, J. "Methods for Generating a Distribution of Optimal Solutions to Nondeterministic Polynomial Optimization Problems", Provisional filed
- 2. Frasch, W.D., Spetzler, D., and York, J. "High Speed, High Fidelity, High Sensitivity Nucleic Acid Detection", Provisional Filed
- 3. Frasch, W.D. and Spetzler, D. "Self-Regulating Answer Formation in DNA Computing via 3D Assembly of DNA", Provisional Filed
- 4. Frasch, W. D. and Xiong, F. "Methods to Use a Double Lock Probe for Rapid and Quantitative Answer Determination of Nondeterministic Polynomial Optimization Problems, Provisional Filed.

Invited Presentations during this Past Year:

- 1. U.S. Army Research Office, Life Sciences Division, Research Triangle Park, NC "DNA Computing"
- 2. Edgewood Chemical Biological Center, U. S. Army RDECOM, Edgewood, MD, "Single-Molecule Detection of DNA via Sequence-Specific Links between F1-ATPase Motors and Gold Nanorod Sensors"
- 3. University of Pittsburgh, Department of Structural Biology, "The F1Fo ATP synthase-two molecular motors with two intertwined mechanisms."
- 4. 13th International Meeting on DNA Computing (DNA13), Memphis, TN, "Probabilistic DNA Computing Solution to a Fully Connected 10-City Traveling Salesman Problem."
- 5. FASEB Summer Research Conference on Transport ATPases: Genomics, Mechanisms, and Relevance to Diseases, Saxtons River, VT "Mechanism of Torque Generation by *E. coli* Biomolecular Motor." F1ATPase
- 6. Gordon Research Conference on Molecular and Cellular Bioenergetics, Andover, NH, "Mechanism of Torque Generation by *E. coli* Biomolecular Motor." F1ATPase
- 7. University of Illinois Urbana Champaign, Department of Biophyics, "Biomolecular motor-powered nanodevices for DNA detection, and their application for DNA Computing."
- 8. University of Illinois Urbana Champaign, Department of Biochemistry, "Single Molecule Studies to Investigate the Rotary Mechanism of the F1Fo molecular motor." 9. University of North Carolina-Chapel Hill, NC, Department of Biochemistry and Biophysics, "Single Molecule Studies to Investigate the Rotary Mechanism of the F1Fo molecular motor."

Awards Past Year:

Best Graduate Student Paper at DNA 13 Conference awarded to David Spetzler, PhD student in my lab.

Scientific or Technological Transitions this Year:

New Funding from Science Foundation Arizona- Small Catalytic Business Grant for Biosensing Project.

Founded Zeptometrics, a new biotechnology company, beginning to build a prototype of a commercial instrument for Biosensing using the technology developed by AFOSR. Initial instrument design will be for rapid detection of *Staphyloccocus aureus* and MRSA. Ultimate goal is to build a hand-held device.

Began hospital trials using the device on patient samples.

